## SEQUENCE LISTING

- <110> TING, ALICE
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- <130> M00656.70097.US
- <140> NOT YET ASSIGNED
- <141> 2003-08-05
- <150> US 60/425,578
- <151> 2002-11-12
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- <170> PatentIn version 3.2
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- <213> Homo sapiens
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Met Glu Ile Lys Leu Glu Asn Lys Tyr Gln Lys Met Glu Asp Phe Ile 50 55 60

Tyr Asp Ala Arg Leu Val Phe Asn Asn Cys Arg Met Tyr Asn Gly Glu 65 70 75 80

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Pro Thr Leu Val Thr Thr Leu Thr Trp Gly Val Gln Cys Phe Ser Arg 65 70 75 80

Tyr Pro Asp His Met Lys Gln His Asp Phe Phe Lys Ser Ala Met Pro 85 90 95

Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn 100 105 110

Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu Gly Asp Thr Leu Val Asn 115 120 125

Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu 130 135 140

Gly His Lys Leu Glu Tyr Asn Tyr Ile Ser His Asn Val Tyr Ile Thr 145 150 155

Ala Asp Lys Gln Lys Asn Gly Ile Lys Ala His Phe Lys Ile Arg His 165 170 175

Asn Ile Glu Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn 180 185 190

Thr Pro Ile Gly Asp Gly Pro Val Leu Leu Pro Asp Asn His Tyr Leu 200 Ser Thr Gln Ser Ala Leu Ser Lys Asp Pro Asn Glu Lys Arg Asp His 215 Met Val Leu Leu Glu Phe Val Thr Ala Ala Arg Met His Lys Arg Gly Pro His Asp Ala Ala Ile Gln Asn Ile Leu Thr Glu Leu Gln Asn His 250 Ala Ala Ala Trp Pro Phe Leu Gln Pro Val Asn Lys Glu Glu Val Pro Asp Tyr Tyr Asp Phe Ile Lys Glu Pro Met Asp Leu Ser Thr Met Glu Ile Lys Leu Glu Ser Asn Lys Tyr Gln Lys Met Glu Asp Phe Ile Tyr Asp Ala Arg Leu Val Phe Asn Asn Cys Arg Met Tyr Asn Gly Glu Asn Thr Ser Tyr Tyr Lys Tyr Ala Asn Arg Leu Glu Lys Phe Phe Asn Asn Lys Val Lys Glu Ile Pro Glu Tyr Ser His Leu Ile Asp Gly Ser Thr 340 345 Ser Gly Ser Gly Lys Pro Gly Ser Gly Glu Gly Ser Ser Gly Arg Gly Lys Gly Gly Lys Gly Leu Gly Lys Gly Gly Ala Lys Arg His Arg Lys Val Leu Arg Asp Asn Ile Gln Gly Ile Thr Glu Leu Met Val Ser Lys 390 395 Gly Glu Glu Leu Phe Thr Gly Val Val Pro Ile Leu Val Glu Leu Asp Gly Asp Val Asn Gly His Lys Phe Ser Val Ser Gly Glu Gly Glu Gly Asp Ala Thr Tyr Gly Lys Leu Thr Leu Lys Phe Ile Cys Thr Thr Gly 440 Lys Leu Pro Val Pro Trp Pro Thr Leu Val Thr Thr Phe Gly Tyr Gly Leu Met Cys Phe Ala Arg Tyr Pro Asp His Met Lys Gln His Asp Phe Phe Lys Ser Ala Met Pro Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe 485 490 Phe Lys Asp Asp Gly Asn Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu 505 Gly Asp Thr Leu Val Asn Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys

520

515

Glu Asp Gly Asn Ile Leu Gly His Lys Leu Glu Tyr Asn Tyr Asn Ser 530 His Asn Val Tyr Ile Met Ala Asp Lys Gln Lys Asn Gly Ile Lys Val Asn Phe Lys Ile Arg His Asn Ile Glu Asp Gly Ser Val Gln Leu Ala Asp His Tyr Gln Gln Asn Thr Pro Ile Gly Asp Gly Pro Val Leu Leu 585 Pro Asp Asn His Tyr Leu Ser Tyr Gln Ser Ala Leu Ser Lys Asp Pro 595 600 Asn Glu Lys Arg Asp His Met Val Leu Leu Glu Phe Val Thr Ala Ala Gly Ile Thr Leu Gly Met Asp Glu Leu Tyr Lys 630 <210> 6 <211> 2412

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Pro Thr Leu Val Thr Thr Leu Thr Trp Gly Val Gln Cys Phe Ser Arg 70 75 80

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Glu Gly Tyr Val Gln Glu Arg Thr Ile Phe Phe Lys Asp Asp Gly Asn 100 105 110

Tyr Lys Thr Arg Ala Glu Val Lys Phe Glu Gly Asp Thr Leu Val Asn 115 120 125

Arg Ile Glu Leu Lys Gly Ile Asp Phe Lys Glu Asp Gly Asn Ile Leu 130 135 140

Gly His Lys Leu Glu Tyr Asn Tyr Ile Ser His Asn Val Tyr Ile Thr 145 150 155 160

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Asn Ala Lys Val Val Lys Asp Tyr Tyr Lys Ile Ile Thr Arg Pro Met 290 295 300

Asp Leu Gln Thr Leu Arg Glu Asn Val Arg Lys Arg Leu Tyr Pro Ser 305 310 315 320

Arg Glu Glu Phe Arg Glu His Leu Glu Leu Ile Val Lys Asn Ser Ala 325 330 335

Thr Tyr Asn Gly Pro Lys His Ser Leu Thr Gln Ile Ser Gln Ser Met 340 345 350

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Ile Gly Asp Gly Pro Val Leu Leu Pro Asp Asn His Tyr Leu Ser Tyr
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